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JC10 REC'D/PU/PD 22 MAR 2002

CLIENT Ref. # BTGI-0025

IN THE UNITED STATES RECEIVING OFFICE
Before the United States Receiving Office
for the Patent Cooperation Treaty

In re application of: Crisanto GUTIERREZ-ARMENTA and Elena RAMIREZ-PARRA,
et al.

U.S. Serial No.:

International Application No.: PCT/EP00/09325

International Filing Date: 25 September 2000 (25.09.2000)

For: WHEAT DP PROTEINS AND USES THEREOF

Assistant Commissioner for Patents
BOX PCT DO/EO/US
Washington, DC 20231

TRANSMITTAL LETTER SUBMITTING SEQUENCE LISTING

Applicant submits herewith the Sequence Listing in written (11 sheets) and computer readable form (1 diskette), the contents of which are the same.

The enclosed Sequence Listing does not go beyond the disclosure in the international application as filed.

Respectfully submitted,



Paul K. Legaard
Registration No. 38,534

/lw

Enc.: 1 Diskette ; 11 sheets sequence listing

Date: 22 March 2002

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SEQUENCE LISTING

<110> Gutierrez-Armenta, Crisanto
Ramirez-Parra, Elena

<120> Wheat DP Proteins And Uses Thereof

<130> BTGI-0025

<150> P9902127
<151> 1999-09-24

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<151> 2000-09-25

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Ser Val Pro Pro Leu Pro Glu Ala Gly Gly Asn Ala Val Gln Arg Lys	
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Gly Ala Val Asp Pro Asp Lys Asp Arg Lys Lys Glu Lys Ala Ala Ala	
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Pro Arg Ile Thr Gly Trp Gly Leu Arg Glu Tyr Ser Lys Ile Val Cys	
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Ile Ala Leu Arg Val Ile Ala Lys Glu Lys Lys Glu Ile Arg Trp Met	
125 130 135	
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Gly Leu Ser Asn Tyr Arg Tyr Glu Lys Ile Lys Lys Leu Glu Glu Val	
140 145 150 155	

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Arg Lys Glu Leu Val Asn Lys Ile Arg Asn Lys Lys Ala Leu Leu Gln	
160 165 170	
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Glu Ile Glu Lys Gln Phe Asp Asp Leu Gln Asn Ile Lys Leu Arg Asn	
175 180 185	
caa aca ctg gaa agc tca gca gag aat gtt aat ggc atc cgc ctt cca	628
Gln Thr Leu Glu Ser Ser Ala Glu Asn Val Asn Gly Ile Arg Leu Pro	
190 195 200	
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Phe Val Leu Val Lys Thr Ser Arg Lys Ala Arg Val Glu Ile Glu Ile	
205 210 215	
tca gat gac tcg aag ttt gcc cat ttc gag ttc aat ggt gca cca ttc	724
Ser Asp Asp Ser Lys Phe Ala His Phe Glu Phe Asn Gly Ala Pro Phe	
220 225 230 235	
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Thr Leu His Asp Asp Leu Ser Ile Leu Glu Gly Val Arg Arg Asn Ser	
240 245 250	
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Ile Gly Arg Ala Gly Arg Ala Thr Leu His	
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35 40 45

Asp Lys Asp Arg Lys Lys Glu Lys Ala Ala Ala Pro Arg Ile Thr Gly
50 55 60

Trp Gly Leu Arg Glu Tyr Ser Lys Ile Val Cys Glu Lys Val Glu Ala
65 70 75 80

Lys Gly Arg Thr Thr Tyr Asn Glu Val Ala Asp Glu Ile Tyr Ser Glu
85 88 90 95

Leu Lys Ser Met Ala His Ile Gly Gln Gly Phe Asp Glu Lys Asn Ile
100 105 110

Arg Arg Arg Val Tyr Asp Ala Phe Asn Val Leu Ile Ala Leu Arg Val
 115 120 125
 Ile Ala Lys Glu Lys Lys Glu Ile Arg Trp Met Gly Leu Ser Asn Tyr
 130 135 140
 Arg Tyr Glu Lys Ile Lys Lys Leu Glu Val Arg Lys Glu Leu Val
 145 150 155 160
 Asn Lys Ile Arg Asn Lys Lys Ala Leu Leu Gln Glu Ile Glu Lys Gln
 165 170 175
 Phe Asp Asp Leu Gln Asn Ile Lys Leu Arg Asn Gln Thr Leu Glu Ser
 180 185 190
 Ser Ala Glu Asn Val Asn Gly Ile Arg Leu Pro Phe Val Leu Val Lys
 195 200 205
 Thr Ser Arg Lys Ala Arg Val Glu Ile Glu Ile Ser Asp Asp Ser Lys
 210 215 220
 Phe Ala His Phe Glu Phe Asn Gly Ala Pro Phe Thr Leu His Asp Asp
 225 230 235 240
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 245 250 255
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ccgggagtcg ggggtcccgcc agcgccgcgat cgcgagatcg ggctt atg tct ggg ggc	177
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ggc agg ccg ccg gct gcg caa aaa atc ctg cag tct ctg cgc ccg ccc	225
Gly Arg Pro Pro Ala Ala Gln Lys Ile Leu Gln Ser Leu Arg Pro Pro	
5 10 15 20	
ccg gtg ttc tcc acg ccg tcg cgg cct ccc ttc gcc tca ccc gac gac	273
Pro Val Phe Ser Thr Pro Ser Arg Pro Pro Phe Ala Ser Pro Asp Asp	
25 30 35	
tac cac cgc ttt cat gcg ccg act acc cct tct gcc act ggc tcc ggc	321
Tyr His Arg Phe His Ala Pro Thr Thr Pro Ser Ala Thr Gly Ser Gly	
40 45 50	

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aag act aag aac aat aaa gct ggg cct cag aca cct acg cca aat gtt Lys Thr Lys Asn Asn Lys Ala Gly Pro Gln Thr Pro Thr Pro Asn Val 120 125 130	561
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 345 350 355
 cat gca aca aac gct aca caa agc agc aaa tca atc aat gtg gaa tat 1281
 His Ala Thr Asn Ala Thr Gln Ser Ser Lys Ser Ile Asn Val Glu Tyr
 360 365 370
 aat att cag cac agg cag aat act cca caa gat cct agt tct tca aat 1329
 Asn Ile Gln His Arg Gln Asn Thr Pro Gln Asp Pro Ser Ser Ser Asn
 375 380 385
 gat tat gga ggg atg aca agg ata atc cct tca gat gtt aat act gat 1377
 Asp Tyr Gly Gly Met Thr Arg Ile Ile Pro Ser Asp Val Asn Thr Asp
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 405 410 415 420
 tgg gaa aca gca cca gaa gtg cag tgg gac acc gct gtg ttt tta cct 1473
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 425 430 435
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 Glu Asp Val Ser Ile Pro His Ala His His Pro Arg Met Gln Val
 440 445 450
 cca agc atg gat caa cca taaggtcatg gcgggtaaaaa cttgacatat 1569
 Pro Ser Met Asp Gln Pro
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 35 40 45

Thr Gly Ser Gly Gly Ile Gly Ser Gly Gly Val Gly Gly Asp Ile Asp
 50 55 60

Glu Gly Leu Val Ile Arg Thr Gln Leu Lys Arg Lys Ala Thr Arg Glu
 65 70 75 80

Glu Asn Asn Ala Ala Glu Ser Ser Asp Cys Met Ile Val Thr Thr Gly
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Val Thr Gly Asn Pro Leu Leu Thr Pro Val Ser Gly Lys Ala Val Lys
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Asn Ser Lys Ser Lys Thr Lys Asn Asn Lys Ala Gly Pro Gln Thr Pro
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Glu Thr Leu Glu Val Gln Lys Arg Arg Ile Tyr Asp Ile Thr Asn Val
180 185 190

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195 200 205

Trp Lys Gly Leu Asp Asp Ser Gly Val Glu Leu Asn Gly Leu Ser
210 215 220

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Asp Glu Arg Ile Ser Asp Met Arg Glu Lys Leu Arg Gly Leu Thr Glu
245 250 255

Asp Glu Asn Ser Gln Arg Trp Leu Tyr Val Thr Glu Asp Asp Ile Lys
260 265 270

Gly Leu Pro Cys Phe Gln Asn Glu Thr Leu Ile Ala Ile Lys Ala Pro
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His Gly Thr Thr Leu Glu Val Pro Asp Pro Asp Glu Ala Gly Asp Tyr
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Leu Gln Arg Arg Tyr Arg Ile Val Leu Arg Ser Thr Leu Gly Pro Ile
305 310 315 320

Asp Val Tyr Leu Val Ser Gln Phe Asp Asp Gly Phe Glu Asn Leu Gly
325 330 335

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Cys Glu Asp Leu His Ala Thr Asn Ala Thr Gln Ser Ser Lys Ser Ile
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Asn Val Glu Tyr Asn Ile Gln His Arg Gln Asn Thr Pro Gln Asp Pro
370 375 380

Ser Ser Ser Asn Asp Tyr Gly Gly Met Thr Arg Ile Ile Pro Ser Asp
385 390 395 400

Val Asn Thr Asp Ala Asp Tyr Trp Leu Leu Thr Glu Gly Asp Val Ser
405 410 415

Ile Thr Asp Met Trp Glu Thr Ala Pro Glu Val Gln Trp Asp Thr Ala
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Thr Ala Ala Leu Asp Leu Thr Gly Val His Ile Leu Glu Ala Ser Ser
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Ala Val Asp Pro
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Trp Gly Leu Arg Glu Tyr Ser Lys Ile Val Cys Glu Asn Val Glu Ala
20 25 30
aaa gga aga aca aca tac aat gag gtt gca gac gaa att tat tca gag 144
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Leu Lys Ser
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Leu Lys Ser
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Val Tyr Asp Ala Phe Asn Val Leu Ile Ala Leu Arg Val Ile Ala Lys

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25

30

gaa aaa aag gag ata cgg tgg atg ggc ctt tca aat tac aga tat gaa
 Glu Lys Lys Glu Ile Arg Trp Met Gly Leu Ser Asn Tyr Arg Tyr Glu
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144

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168

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Lys Ile Lys Lys Leu Glu Glu Val
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48

gaa atc gaa aaa cag ttt gat gat ctc caa aac atc aag tta cgt aac
 Glu Ile Glu Lys Gln Phe Asp Asp Leu Gln Asn Ile Lys Leu Arg Asn
 20 25 30

96

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Phe Val Leu Val Lys Thr Ser Arg
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Phe Glu Phe Asn Gly Ala Pro Phe Thr Leu His Asp Asp Leu Ser Ile
20 25 30

ctt gag ggg gta agg cgt aac agc ata gga aga gct ggc cgc gcc acc 144
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35 40 45

ctt cac 150
Leu His
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<213> Triticum monococcum

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Phe Glu Phe Asn Gly Ala Pro Phe Thr Leu His Asp Asp Leu Ser Ile
20 25 30

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35 40 45

Leu His
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<213> Artificial Sequence

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